

# **A novel motif in the V3 domain of PKC- $\theta$ determines its immunological synapse localization and functions in T cells via association with CD28**

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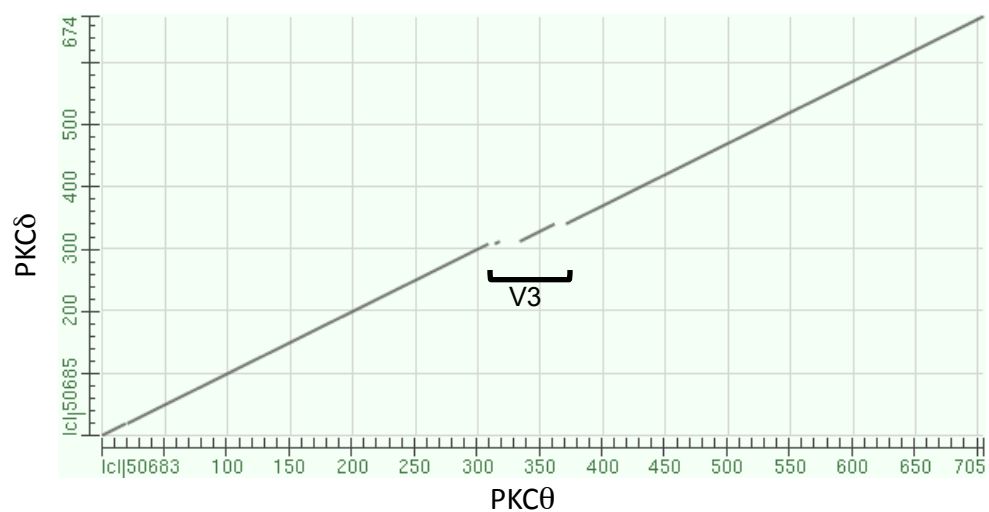
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Supporting Online Materials

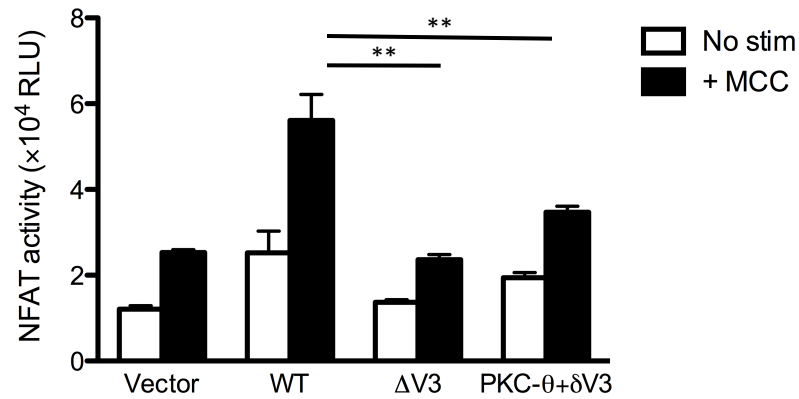
Supplementary Figures and Legends 1 – 9

**Figure S1**



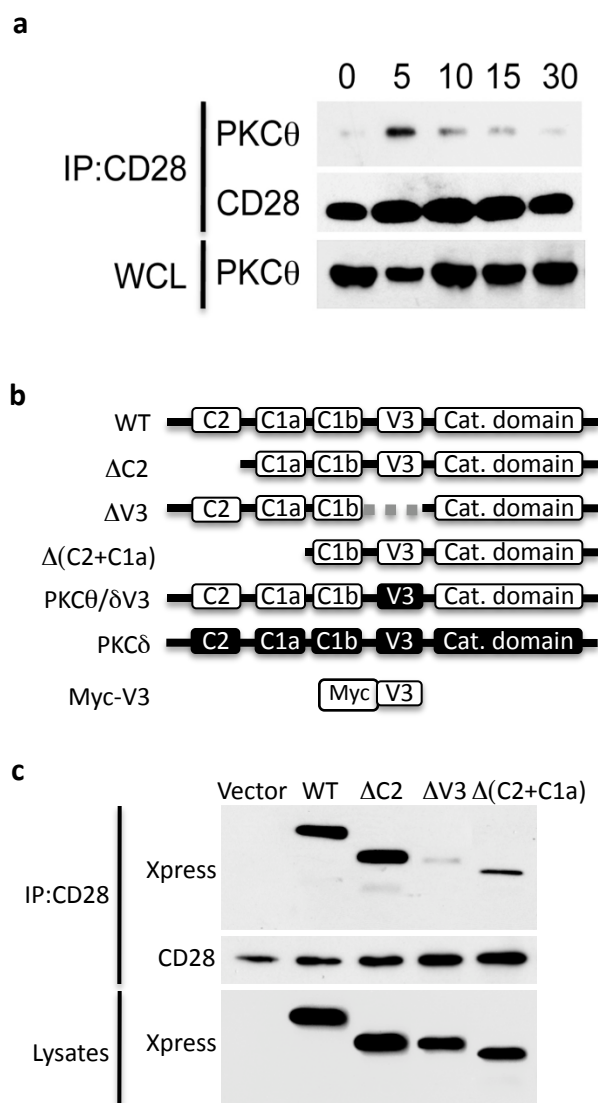
**Supplementary Figure 1.** Alignment of human PKC- $\theta$  (NP\_006248) and PKC- $\delta$  (NP\_006245) showing the divergence of their V3 (hinge) regions. Sequences were aligned using the NCBI BLAST program.

**Figure S2**



**Supplementary Figure 2.** Importance of the PKC-θ V3 domain for NFAT activation. Normalized Luc activity in MCC-specific T hybridoma cells cotransfected with empty pEF vector, WT PKC-θ, PKC-θ-ΔV3, or PKC-θ+δV3 together with NFAT-Luc and β-Gal reporter plasmids. Cells were cultured with DCEK fibroblasts expressing I-E<sup>k</sup> and B7-1 in the absence or presence of MCC peptide for 6 h. \*\* p < .05. Data are from two experiments.

**Figure S3**



**Supplementary Figure 3.** The PKC-θ V3 domain interacts with CD28. (a) Immunoblot analysis of CD28 IPs or whole cell lysates (WCL) from Jurkat E6.1 T cells left unstimulated or stimulated for the indicated times (min) with anti-CD3 plus anti-CD28 mAbs. (b) Schematic representation of PKC-θ mutants. (c) Immunoblot analysis of CD28 IPs or lysates from Jurkat E6.1 T cells transfected with an empty vector or the indicated PKC-θ vectors, and left unstimulated or stimulated for 5 min with anti-CD3 plus anti-CD28 mAbs.

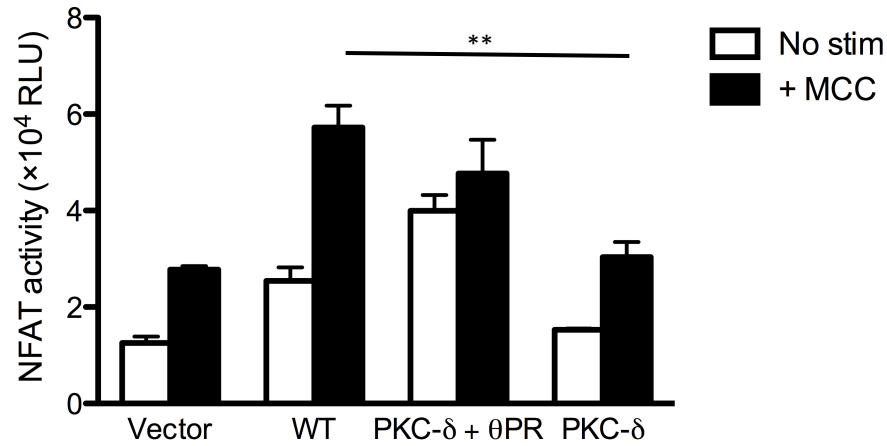
**Figure S4**



Species	Sequence	Ensembl Accession
<i>Homo sapiens/Gorilla gorilla</i>	AR <u>P</u> <u>P</u> CL <u>P</u> TP	ENST00000263125
<i>Pan troglodytes</i>	AR <u>P</u> <u>P</u> CL <u>P</u> TL	ENSPTRP00000041216
<i>Macaca mulatta</i>	AR <u>P</u> <u>P</u> CL <u>P</u> TP	ENSMMPUP00000027347
<i>Canis familiaris</i>	ARL <u>P</u> CV <u>P</u> AP	ENSCAFP00000007725
<i>Felis catus</i>	ARL <u>P</u> CV <u>P</u> AS	ENSFCAP00000008789
<i>Equus caballus</i>	AKL <u>P</u> HAP <u>P</u> AP	ENSECAP00000020818
<i>Bos taurus</i>	AKP <u>P</u> YV <u>P</u> GP	ENSBTAT00000060978
<i>Loxodonta africana</i>	TRL <u>P</u> YL <u>P</u> TP	ENSLAFP00000001356
<i>Ailuropoda melanoleuca</i>	AKL <u>P</u> CV <u>P</u> AP	EFB18582.1 (NCBI)
<i>Mus musculus</i>	TRP <u>P</u> CV <u>P</u> TP	ENSMUST00000028118
<i>Rattus norvegicus</i>	TRP <u>P</u> CV <u>P</u> TP	ENSRNOP00000025902
<i>Ochotona princeps</i>	TRP <u>P</u> YL <u>P</u> TP	ENSOPRP00000002826
<i>Dipodomys ordii</i>	TRQ <u>P</u> NF <u>P</u> TP	ENSDORP00000014980
<i>Spermophilus tridecemlineatus</i>	AR <u>P</u> YL <u>P</u> TP	ENSSTOP00000008114
<i>Tupaia belangeri</i>	ARS <u>P</u> YL <u>P</u> TP	ENSTBEP00000011279
<i>Procapra capensis</i>	TRL <u>P</u> YL <u>P</u> TP	ENSPCAP00000013723
<i>Echinops telfairi</i>	TKL <u>P</u> YL <u>P</u> AP	ENSETEP00000013887
<i>Cavia porcellus</i>	ARL <u>P</u> YL <u>P</u> TG	ENSCPOP00000013395
<i>Dasyus novemcinctus</i>	TRL <u>P</u> YL <u>P</u> VP	ENSDNOP00000008621
<i>Pteropus vampyrus</i>	AR <u>P</u> PHG <u>P</u> AL	ENSPVAP00000014575
<i>Tursiops truncatus</i>	AKL <u>P</u> YG <u>P</u> AP	ENSTTRP00000012525
<i>Xenopus laevis</i>	PKA <u>P</u> GL <u>P</u> MP	BAC79120.1 (NCBI)
<i>Danio rerio</i>	AIS <u>P</u> LT <u>P</u> AP	ENSDART00000046253
<i>Tetraodon nigroviridis</i>	LLL <u>P</u> NL <u>P</u> LP	CAG04125.1 (NCBI)
<i>Takifugu rubripes</i>	VR <u>P</u> SG <u>P</u> IT	ENSTRUP00000030203

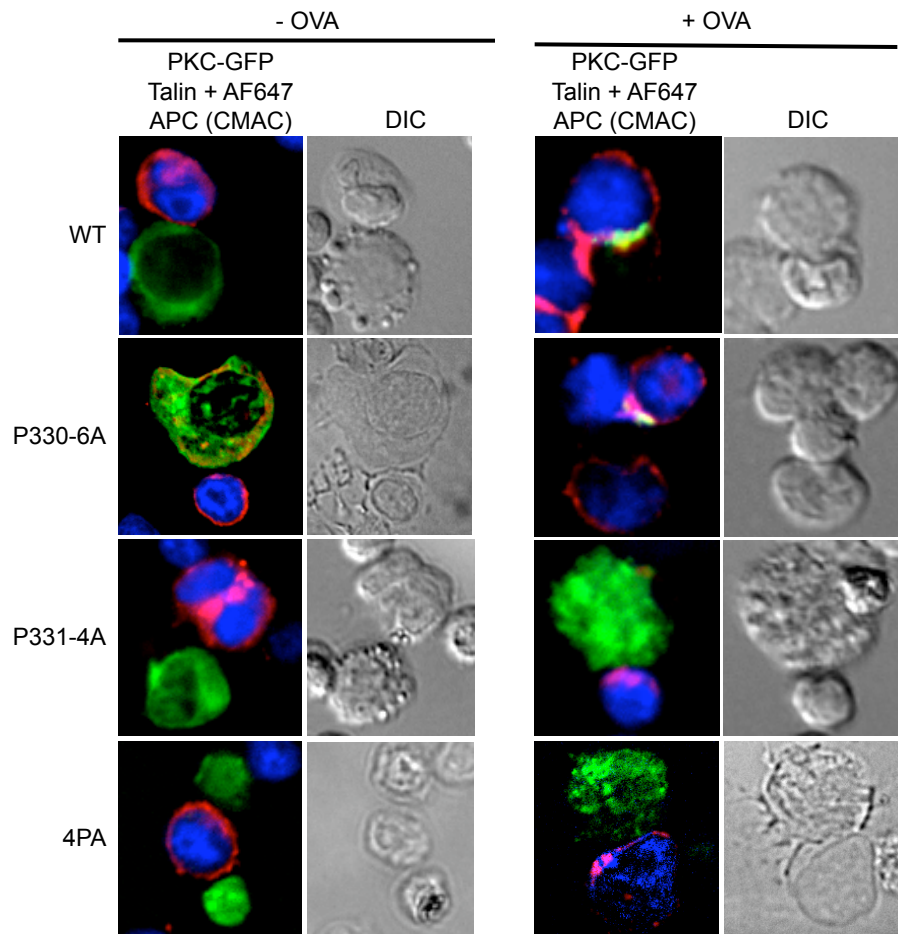
**Supplementary Figure 4.** Evolutionary conservation of the PxxP motif in the V3 domain of PKC-θ. Sequence alignment of the PR motif in putative PKC-θ enzymes from the indicated organisms. The human sequence corresponds to amino acid 328-336 of human PKC-θ. Alignment was performed using Weblogo (<http://weblogo.berkeley.edu/logo.cgi>). Proline residues that are absolutely conserved in all species are underlined in bold.

**Figure S5**



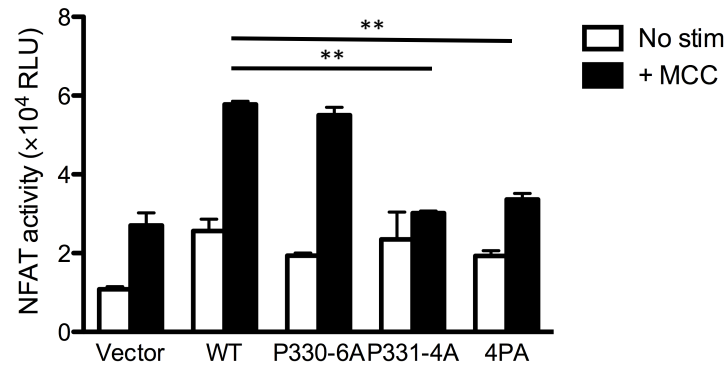
**Supplementary Figure 5.** Importance of the PR motif in the V3 domain of PKC-θ for NFAT activation. Normalized Luc activity in MCC-specific T hybridoma cells cotransfected with empty pEF vector or the indicated PKC-θ vectors together with NFAT-Luc and β-Gal reporter plasmids. Cells were cultured with DCEK fibroblasts expressing I-E<sup>k</sup> and B7-1 in the absence or presence of MCC peptide for 6 h. \*\* p < .05. Data are from two experiments.

**Figure S6**



**Supplementary Fig. 6.** Importance of the PxxP motif in the V3 domain of PKC- $\theta$  for IS localization and CD28 interaction. *PKC $\theta$ <sup>-/-</sup>* OT-II CD4<sup>+</sup> T cells were infected with retrovirus expressing GFP-tagged PKC- $\theta$ , or PKC  $\theta$ -GFP fusion vectors containing mutations at P330-6A, P331-4A, or all four proline residues (4PA) (green). Cells were fixed, stained and analyzed as in **Fig. 1a**.

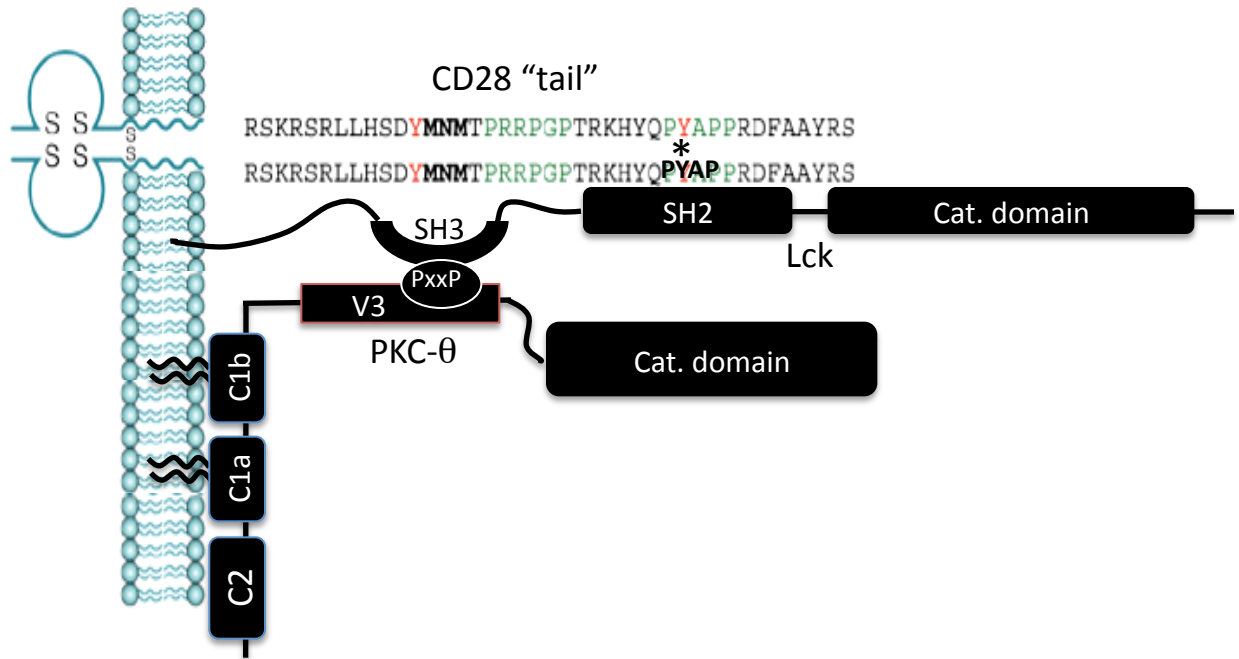
**Figure S7**



**Supplementary Figure 7.** A PxxP motif in V3 affects PKC- $\theta$ -mediated NFAT signaling. Normalized Luc activity in MCC-specific T hybridoma cells cotransfected with empty pEF vector or the indicated PKC- $\theta$  vectors together with NFAT-Luc and  $\beta$ -Gal reporter plasmids. Cells were cultured with DCEK fibroblasts expressing I-E<sup>k</sup> and B7-1 in the absence or presence of MCC peptide for 6 h. \*\* p < .05. Data are from two experiments.

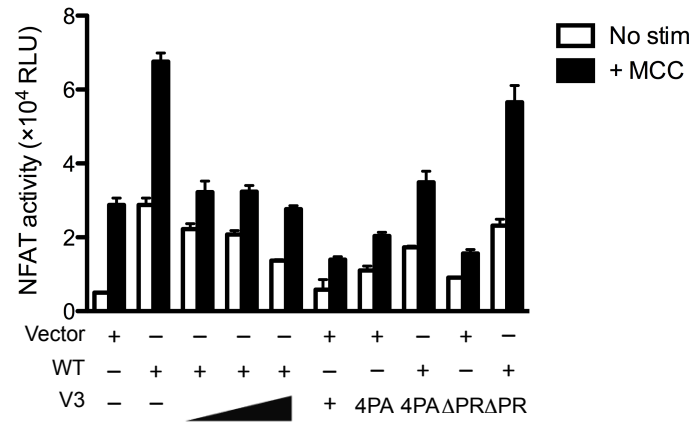


**Figure S8**



**Supplementary Figure 8.** Schematic mode of tri-partite interaction between CD28, Lck and PKC-θ. Lck mediates the interaction between CD28 and PKC-θ, with its SH3 domain binding the PR motif in the V3 domain of PKC-θ, and its SH2 domain binding phosphorylated Tyr-207 in the CD28 P<sup>206</sup>Y\*AP<sup>209</sup> motif.

**Figure S9**



**Supplementary Figure 9.** The PKC- $\theta$  V3 domain interferes with PKC- $\theta$ -mediated NFAT activation. Normalized Luc activity in MCC-specific T hybridoma cells cotransfected with empty pEF vector or the indicated combinations of full-length PKC $\theta$  and/or PKC $\theta$ -V3 vectors together with NFAT-Luc and  $\beta$ -Gal reporter plasmids. Cells were cultured with DCEK fibroblasts expressing I-E<sup>k</sup> and B7-1 in the absence or presence of MCC peptide for 6 h. Data are from two experiments.